SEQUENCE LISTING

<110>  Haskell-Luevano, Carrie

<120>  Novel Melanocortin Receptor Peptide Template for the Treatment of
       Obesity

<130>  UF-375

<160>  43

<170>  PatentIn version 3.2

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       (begin)

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Tyr Lys Pro Val
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<223>  Xaa = diaminoproprionic acid (Dpr)


<220>
<221>  MISC_FEATURE
<222>  (12)..(12)
<223>  Xaa in this location represents the cyclization of this peptide
       (end)


<400>  33

Tyr Xaa Asp His Xaa Arg Trp Asn Ala Phe Xaa Xaa Tyr
1               5                   10



<210>  34
<211>  13
<212>  PRT
<213>  Artificial sequence


<220>
<223>  chimeric peptide



<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
<221>  MISC_FEATURE
<222>  (5)..(5)
<223>  Xaa = DNal(1')


<220>
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<223>  Xaa = diaminoproprionic acid (Dpr)


<220>
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<222>  (12)..(12)
<223>  Xaa in this location represents the cyclization of this peptide
       (end)


<400>  34

Tyr Xaa Asp His Xaa Arg Trp Asn Ala Phe Xaa Xaa Tyr
1               5                   10



<210>  35
<211>  13
<212>  PRT
<213>  Artificial sequence
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<220>
<223>  chimeric peptide


<220>
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<222>  (2)..(2)
<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
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<222>  (5)..(5)
<223>  Xaa = DBip


<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (end)


<400>  35

Tyr Xaa Asp His Xaa Arg Trp Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  36
<211>  13
<212>  PRT
<213>  Artificial Sequence


<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
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<223>  Xaa = DPhe


<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (end)

<400>  36

Tyr Xaa Asp His Xaa Pro Trp Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  37
<211>  13
<212>  PRT
<213>  Artificial sequence

<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)

<220>
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<222>  (5)..(5)
<223>  Xaa = DPhe

<220>
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<220>
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<222>  (12)..(12)
<223>  Xaa in this location represents the cyclization of this peptide
       (end)

<400>  37

Tyr Xaa Asp His Xaa Lys Trp Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  38
<211>  13
<212>  PRT
<213>  Artificial sequence

<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
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<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (end)


<400>  38

Tyr Xaa Asp His Xaa Arg Pro Asn Ala Phe Xaa Xaa Tyr
1               5                   10



<210>  39
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<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
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<223>  Xaa = DPhe


<220>
<221>  MISC_FEATURE
<222>  (7)..(7)
<223>  Xaa = Nal(2')


<220>
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<223>  Xaa = diaminoproprionic acid (Dpr)

<220>
<221>  MISC_FEATURE
<222>  (12)..(12)
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       (end)

<400>  39

Tyr Xaa Asp His Xaa Arg Xaa Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  40
<211>  13
<212>  PRT
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<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)

<220>
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<223>  Xaa = DPhe

<220>
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<223>  Xaa = DNal(2')

<220>
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<220>
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<400>  40

Tyr Xaa Asp His Xaa Arg Xaa Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  41
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<211>  13
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<220>
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<220>
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<223>  Xaa in this location represents the cyclization of this peptide
       (begin)

<220>
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<220>
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<220>
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<400>  41

Tyr Xaa Asp His Xaa Arg Xaa Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  42
<211>  13
<212>  PRT
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<220>
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<220>
<221>  MISC_FEATURE
<222>  (2)..(2)
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       (begin)

<220>
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<221>  MISC_FEATURE
<222>  (5)..(5)
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<220>
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<220>
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<220>
<221>  MISC_FEATURE
<222>  (12)..(12)
<223>  Xaa in this location represents the cyclization of this peptide
       (end)


<400>  42

Tyr Xaa Asp His Xaa Arg Xaa Asn Ala Phe Xaa Xaa Tyr
1               5                   10


<210>  43
<211>  13
<212>  PRT
<213>  Artificial sequence


<220>
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<220>
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<222>  (2)..(2)
<223>  Xaa in this location represents the cyclization of this peptide
       (begin)


<220>
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<220>
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<220>
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<221> MISC_FEATURE
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<223> Xaa in this location represents the cyclization of this peptide
      (end)

<400> 43

Tyr Xaa Asp His Xaa Arg Trp Asn Ala Xaa Xaa Xaa Tyr
1               5                   10